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1641

TECH CENTER 1600/2900

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/756,899A

DATE: 05/03/2001

TIME: 15:59:47

Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\05032001\I756899A.raw

ENTERED

3 <110> APPLICANT: Redgeld, Franciscus
 4 Kraneveld, Aletta
 5 Nijkamp, Franciscus
 7 <120> TITLE OF INVENTION: INHIBITION OF PROTEIN BINDING TO MAST CELLS
 9 <130> FILE REFERENCE: 2183-4692
 11 <140> CURRENT APPLICATION NUMBER: US 09/756,899A
 12 <141> CURRENT FILING DATE: 2001-01-09
 14 <150> PRIOR APPLICATION NUMBER: PCT/NL99/00430
 15 <151> PRIOR FILING DATE: 1997-07-07
 17 <160> NUMBER OF SEQ ID NOS: 3
 19 <170> SOFTWARE: PatentIn version 3.0
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 9
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Unknown
 26 <220> FEATURE:
 27 <223> OTHER INFORMATION: peptide
 29 <400> SEQUENCE: 1
 31 Ala His Trp Ser Gly His Cys Cys Leu
 32 1 5
 34 <210> SEQ ID NO: 2
 35 <211> LENGTH: 16
 36 <212> TYPE: PRT
 37 <213> ORGANISM: Unknown
 39 <220> FEATURE:
 40 <223> OTHER INFORMATION: protein with apparent molecular weight of 27kDal
 42 <220> FEATURE:
 43 <221> NAME/KEY: misc_feature
 44 <222> LOCATION: (1)..(16)
 45 <223> OTHER INFORMATION: Xaa can be any amino acid
 48 <400> SEQUENCE: 2
 50 Asp Ile Gln Met Thr Gln Ser Pro Pro Ser Leu Ser Ala Xaa Leu Gly
 51 1 5 10 15
 53 <210> SEQ ID NO: 3
 54 <211> LENGTH: 16
 55 <212> TYPE: PRT
 56 <213> ORGANISM: Unknown
 58 <220> FEATURE:
 59 <223> OTHER INFORMATION: N-terminal amino acid sequence
 61 <400> SEQUENCE: 3
 63 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly
 64 1 5 10 15

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/756,899A

DATE: 05/03/2001

TIME: 15:59:48

Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\05032001\I756899A.raw

L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

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Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/756,899

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 8 Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of <213>Organism (NEW RULES) Sequence(s) are missing this mandatory field or its response.
- 12 Use of <220>Feature (NEW RULES) Sequence(s) 1-2 are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted "file," resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/756,899

DATE: 01/25/2001
 TIME: 10:43:14

Input Set : A:\sequence listing.txt
 Output Set: N:\CRF3\01252001\I756899.raw

Does Not Comply
 Corrected Diskette Needed

3 <110> APPLICANT: Redegeld, Franciscus
 4 Kraneveld, Aletta
 5 Nijkamp, Franciscus
 7 <120> TITLE OF INVENTION: INHIBITION OF PROTEIN BINDING TO MAST CELLS
 9 <130> FILE REFERENCE: 4692 US

OK> 11 <140> CURRENT APPLICATION NUMBER: US/09/756,899
 11 <141> CURRENT FILING DATE: 2001-01-09

11 <150> PRIOR APPLICATION NUMBER: PCT/NL99/00430,
 12 <151> PRIOR FILING DATE: 1999-07-07

14 <160> NUMBER OF SEQ ID NOS: 3

16 <170> SOFTWARE: PatentIn version 3.0

18 <210> SEQ ID NO: 1

19 <211> LENGTH: 9

20 <212> TYPE: PRT

21 <213> ORGANISM: unidentified

23 <400> SEQUENCE: 1

25 Ala His Trp Ser Gly His Cys Cys Leu

26 1 5

28 <210> SEQ ID NO: 2

29 <211> LENGTH: 16

30 <212> TYPE: PRT

31 <213> ORGANISM: unidentified

33 <220> FEATURE:

34 <221> NAME/KEY: misc_feature

35 <223> OTHER INFORMATION: Xaa can be any amino acid

38 <400> SEQUENCE: 2

OK> 40 Asp Ile Gln Met Thr Gln Ser Pro Pro Ser Leu Ser Ala Xaa Leu Gly

41 1 5 10 15

43 <210> SEQ ID NO: 3

44 <211> LENGTH: 16

45 <212> TYPE: PRT

46 <213> ORGANISM: Immunoglobulin LC

48 <400> SEQUENCE: 3

50 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly

51 1 5 10 15

Per 1.823 of new Sequence Rules, the only valid
 <213> responses are: Unknown, Artificial Sequence,
 or scientific name

see item 12
 on Error Summary
 sheet

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/756,899

DATE: 01/25/2001

TIME: 10:43:15

Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\01252001\I756899.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:40 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:2
L:40 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2